

This page gives you Search Results detail for the Application 10578613 and Search Result 20090820_143901_us-10-578-613-1_copy_1_659.rge.

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GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: August 22, 2009, 17:09:10 ; Search time 628 Seconds
(without alignments)
115985.772 Million cell updates/sec

Title: US-10-578-613-1_COPY_1_659
Perfect score: 659
Sequence: 1 gatttgccgatgcaacaggc.....tatttcggcagctcttgca 659

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 18669703 seqs, 55253738715 residues

Total number of hits satisfying chosen parameters: 37339406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*
1: gb_bct:*
2: gb_env:*
3: gb_htc:*
4: gb_htgl:*
5: gb_htg2:*
6: gb_htg3:*
7: gb_inv:*
8: gb_mam:*
9: gb_pat1:*
10: gb_pat2:*
11: gb_phg:*
12: gb_pln:*
13: gb_pri:*
14: gb_rod:*
15: gb_sts:*
16: gb_syn:*
17: gb_tsa:*
18: gb_una:*
19: gb_vrl:*
20: gb_vrt:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	659	100.0	662	9	DD081529	DD081529 Recombina
2	659	100.0	3150	1	AB018420	AB018420 Bacillus
3	659	100.0	3150	9	BD186021	BD186021 Host micr
4	659	100.0	3150	9	BD277536	BD277536 Host micr
5	659	100.0	3150	9	CS097336	CS097336 Sequence
6	659	100.0	3150	9	CS144748	CS144748 Sequence
7	659	100.0	3150	9	DD017832	DD017832 Host micr
8	659	100.0	3150	9	DD170905	DD170905 Host micr
9	659	100.0	3150	9	DD170944	DD170944 Host micr
10	659	100.0	3150	9	DD172235	DD172235 Recombina
11	659	100.0	3150	9	DD226462	DD226462 Modified
12	659	100.0	3150	9	DD226988	DD226988 Mutant Ba
13	659	100.0	3150	9	DD235791	DD235791 Modified
14	659	100.0	3150	9	DD235806	DD235806 Mutant Ba
15	659	100.0	3150	9	DD259153	DD259153 Host micr
16	659	100.0	3150	9	DD367975	DD367975 Recombina
17	659	100.0	3150	9	DD378746	DD378746 Recombina
18	659	100.0	3150	9	DD401230	DD401230 Host micr
19	659	100.0	3150	9	DD434758	DD434758 host micr
20	659	100.0	3150	9	DD434783	DD434783 Recombina
21	659	100.0	3150	9	DD452572	DD452572 Recombina
22	659	100.0	3150	9	DD452652	DD452652 Recombina
23	659	100.0	3150	9	DD460426	DD460426 Recombina
24	659	100.0	3150	9	DD490779	DD490779 Recombina
25	659	100.0	3150	9	DD490874	DD490874 Recombina
26	659	100.0	3150	9	DD491790	DD491790 Recombina
27	659	100.0	3150	9	DD495339	DD495339 Recombina
28	659	100.0	3150	9	DD495383	DD495383 Host micr
29	659	100.0	3150	9	DJ029919	DJ029919 Novel var
30	659	100.0	3150	9	DJ046641	DJ046641 Recombina
31	659	100.0	3150	9	DJ046789	DJ046789 Novel var
32	659	100.0	3150	9	DJ085180	DJ085180 Recombina
33	659	100.0	3150	9	DJ400174	DJ400174 Recombina
34	659	100.0	3150	9	DJ400225	DJ400225 Recombina
35	659	100.0	3150	9	DJ400270	DJ400270 Recombina
36	659	100.0	3150	9	DJ400333	DJ400333 Recombina
37	659	100.0	3150	9	DJ430230	DJ430230 Recombina
38	659	100.0	3150	9	DL184266	DL184266 Recombina
39	659	100.0	3150	9	DL239996	DL239996 Recombina
40	659	100.0	3150	9	DM015068	DM015068 Recombina
41	659	100.0	3150	9	DM039094	DM039094 Recombina
42	659	100.0	3150	10	GM864698	GM864698 Sequence
43	659	100.0	3189	9	DD320520	DD320520 recombina
44	659	100.0	3189	9	DD320533	DD320533 recombina
45	659	100.0	3189	9	E37675	E37675 Thermostabl